

#5

1/20

cca gcc cta cca gaa gat ggg ggg tcc ggg gcc ttc cca cca ggg cac	48
Pro Ala Leu Pro Glu Asp Gly Gly Ser Gly Ala Phe Pro Pro Gly His	
1 5 10 15	
ttc aaa gat cca aaa cga cta tat tgt aaa aac ggg ggg ttc ttc cta	96
Phe Lys Asp Pro Lys Arg Leu Tyr Cys Lys Asn Gly Gly Phe Phe Leu	
20 25 30	
cga atc cac cca gat ggg cga gta gat ggg gta cga gaa aaa tcc gat	144
Arg Ile His Pro Asp Gly Arg Val Asp Gly Val Arg Glu Lys Ser Asp	
35 40 45	
cca cac atc aaa cta caa cta caa gcc gaa gaa cga ggg gta gta tcc	192
Pro His Ile Lys Leu Gln Leu Gln Ala Glu Glu Arg Gly Val Val Ser	
50 55 60	
atc aaa ggg gta tgt gcc aac cga tat cta gcc atg aaa gaa gat ggg	240
Ile Lys Gly Val Cys Ala Asn Arg Tyr Leu Ala Met Lys Glu Asp Gly	
65 70 75 80	
cga cta cta gcc tcc aaa tgt gta acc gat gaa tgt ttc ttc ttc gaa	288
Arg Leu Leu Ala Ser Lys Cys Val Thr Asp Glu Cys Phe Phe Phe Glu	
85 90 95	
cga cta gaa tcc aac aac tat aac acc tat cga tcc cga aaa tat tcc	336
Arg Leu Glu Ser Asn Asn Tyr Asn Thr Tyr Arg Ser Arg Lys Tyr Ser	
100 105 110	
tcc tgg tat gta gcc cta aaa cga acc ggg caa tat aaa cta ggg cca	384
Ser Trp Tyr Val Ala Leu Lys Arg Thr Gly Gln Tyr Lys Leu Gly Pro	
115 120 125	
aaa acc ggg cca ggg caa aaa gcc atc cta ttc cta cca atg tcc gcc	432
Lys Thr Gly Pro Gly Gln Lys Ala Ile Leu Phe Leu Pro Met Ser Ala	
130 135 140	
aaa tcc taa	441
Lys Ser *	
145	

FIG. 1.

09886856.100504

2/20

Pro Ala Leu Pro Glu Asp Gly Gly Ser Gly Ala Phe Pro Pro Gly His
1 5 10 15
Phe Lys Asp Pro Lys Arg Leu Tyr Cys Lys Asn Gly Gly Phe Phe Leu
20 25 30
Arg Ile His Pro Asp Gly Arg Val Asp Gly Val Arg Glu Lys Ser Asp
35 40 45
Pro His Ile Lys Leu Gln Leu Gln Ala Glu Glu Arg Gly Val Val Ser
50 55 60
Ile Lys Gly Val Cys Ala Asn Arg Tyr Leu Ala Met Lys Glu Asp Gly
65 70 75 80
Arg Leu Leu Ala Ser Lys Cys Val Thr Asp Glu Cys Phe Phe Phe Glu
85 90 95
Arg Leu Glu Ser Asn Asn Tyr Asn Thr Tyr Arg Ser Arg Lys Tyr Ser
100 105 110
Ser Trp Tyr Val Ala Leu Lys Arg Thr Gly Gln Tyr Lys Leu Gly Pro
115 120 125
Lys Thr Gly Pro Gly Gln Lys Ala Ile Leu Phe Leu Pro Met Ser Ala
130 135 140
Lys Ser
145

FIG. 2.

09886856-100501

3/20

ccc gcc ttg ccc gag gat ggc ggc agc ggc gcc ttc ccg ccc ggc cac	48
Pro Ala Leu Pro Glu Asp Gly Gly Ser Gly Ala Phe Pro Pro Gly His	
1 5 10 15	
ttc aag gac ccc aag cgg ctg tac tgc aaa aac ggg ggc ttc ttc ctg	96
Phe Lys Asp Pro Lys Arg Leu Tyr Cys Lys Asn Gly Gly Phe Phe Leu	
20 25 30	
cgc atc cac ccc gac ggc cga gtt gac ggg gtc cgg gag aag agc gac	144
Arg Ile His Pro Asp Gly Arg Val Asp Gly Val Arg Glu Lys Ser Asp	
35 40 45	
cct cac atc aag cta caa ctt caa gca gaa gag aga gga gtt gtg tct	192
Pro His Ile Lys Leu Gln Leu Gln Ala Glu Glu Arg Gly Val Val Ser	
50 55 60	
atc aaa gga gtg tgt gct aac cgt tac ctg gct atg aag gaa gat gga	240
Ile Lys Gly Val Cys Ala Asn Arg Tyr Leu Ala Met Lys Glu Asp Gly	
65 70 75 80	
aga tta ctg gct tct aaa tgt gtt acg gat gag tgt ttc ttt ttt gaa	288
Arg Leu Leu Ala Ser Lys Cys Val Thr Asp Glu Cys Phe Phe Phe Glu	
85 90 95	
cga ttg gaa tct aat aac tac aat act tac cgg tca agg aaa tac acc	336
Arg Leu Glu Ser Asn Asn Tyr Asn Thr Tyr Arg Ser Arg Lys Tyr Thr	
100 105 110	
agt tgg tat gtg gca ctg aaa cga act ggg cag tat aaa ctt gga tcc	384
Ser Trp Tyr Val Ala Leu Lys Arg Thr Gly Gln Tyr Lys Leu Gly Ser	
115 120 125	
aaa aca gga cct ggg cag aaa gct ata ctt ttt ctt cca atg tct gct	432
Lys Thr Gly Pro Gly Gln Lys Ala Ile Leu Phe Leu Pro Met Ser Ala	
130 135 140	
aag agc tga	441
Lys Ser *	
145	

FIG. 3.

09886856.100501

4/20

atg gca gcc ggg agc atc acc acg ctg cca gcc cta cca gaa gat ggg	48
Met Ala Ala Gly Ser Ile Thr Thr Leu Pro Ala Leu Pro Glu Asp Gly	
1 5 10 15	
ggg tcc ggg gcc ttc cca cca ggg cac ttc aaa gat cca aaa cga cta	96
Gly Ser Gly Ala Phe Pro Pro Gly His Phe Lys Asp Pro Lys Arg Leu	
20 25 30	
tat tgt aaa aac ggg ggg ttc ttc cta cga atc cac cca gat ggg cga	144
Tyr Cys Lys Asn Gly Gly Phe Phe Leu Arg Ile His Pro Asp Gly Arg	
35 40 45	
gta gat ggg gta cga gaa aaa tcc gat cca cac atc aaa cta caa cta	192
Val Asp Gly Val Arg Glu Lys Ser Asp Pro His Ile Lys Leu Gln Leu	
50 55 60	
caa gcc gaa gaa cga ggg gta gta tcc atc aaa ggg gta tgt gcc aac	240
Gln Ala Glu Glu Arg Gly Val Val Ser Ile Lys Gly Val Cys Ala Asn	
65 70 75 80	
cga tat cta gcc atg aaa gaa gat ggg cga cta cta gcc tcc aaa tgt	288
Arg Tyr Leu Ala Met Lys Glu Asp Gly Arg Leu Leu Ala Ser Lys Cys	
85 90 95	
gta acc gat gaa tgt ttc ttc ttc gaa cga cta gaa tcc aac aac tat	336
Val Thr Asp Glu Cys Phe Phe Phe Glu Arg Leu Glu Ser Asn Asn Tyr	
100 105 110	
aac acc tat cga tcc cga aaa tat tcc tcc tgg tat gta gcc cta aaa	384
Asn Thr Tyr Arg Ser Arg Lys Tyr Ser Ser Trp Tyr Val Ala Leu Lys	
115 120 125	
cga acc ggg caa tat aaa cta ggg cca aaa acc ggg cca ggg caa aaa	432
Arg Thr Gly Gln Tyr Lys Leu Gly Pro Lys Thr Gly Pro Gly Gln Lys	
130 135 140	
gcc atc cta ttc cta cca atg tcc gcc aaa tcc taa	468
Ala Ile Leu Phe Leu Pro Met Ser Ala Lys Ser *	
145 150 155	

FIG. 4.

09886856-100501

5/20

atg gca gcc ggg agc atc acc acg ctg ccc gcc ttg ccc gag gat ggc 48
 Met Ala Ala Gly Ser Ile Thr Thr Leu Pro Ala Leu Pro Glu Asp Gly
 1 5 10 15

ggc agc ggc gcc ttc ccg ccc ggc cac ttc aag gac ccc aag cgg ctg 96
 Gly Ser Gly Ala Phe Pro Pro Gly His Phe Lys Asp Pro Lys Arg Leu
 20 25 30

tac tgc aaa aac ggg ggc ttc ttc ctg cgc atc cac ccc gac ggc cga 144
 Tyr Cys Lys Asn Gly Gly Phe Phe Leu Arg Ile His Pro Asp Gly Arg
 35 40 45

gtt gac ggg gtc cgg gag aag agc gac cct cac atc aag cta caa ctt 192
 Val Asp Gly Val Arg Glu Lys Ser Asp Pro His Ile Lys Leu Gln Leu
 50 55 60

caa gca gaa gag aga gga gtt gtg tct atc aaa gga gtg tgt gct aac 240
 Gln Ala Glu Glu Arg Gly Val Val Ser Ile Lys Gly Val Cys Ala Asn
 65 70 75 80

cgt tac ctg gct atg aag gaa gat gga aga tta ctg gct tct aaa tgt 288
 Arg Tyr Leu Ala Met Lys Glu Asp Gly Arg Leu Leu Ala Ser Lys Cys
 85 90 95

gtt acg gat gag tgt ttc ttt ttt gaa cga ttg gaa tct aat aac tac 336
 Val Thr Asp Glu Cys Phe Phe Phe Glu Arg Leu Glu Ser Asn Asn Tyr
 100 105 110

aat act tac cgg tca agg aaa tac acc agt tgg tat gtg gca ctg aaa 384
 Asn Thr Tyr Arg Ser Arg Lys Tyr Thr Ser Trp Tyr Val Ala Leu Lys
 115 120 125

cga act ggg cag tat aaa ctt gga tcc aaa aca gga cct ggg cag aaa 432
 Arg Thr Gly Gln Tyr Lys Leu Gly Ser Lys Thr Gly Pro Gly Gln Lys
 130 135 140

gct ata ctt ttt ctt cca atg tct gct aag agc tga ttttaa 474
 Ala Ile Leu Phe Leu Pro Met Ser Ala Lys Ser *
 145 150 155

FIG. 5.

FOOTNOTES 100501

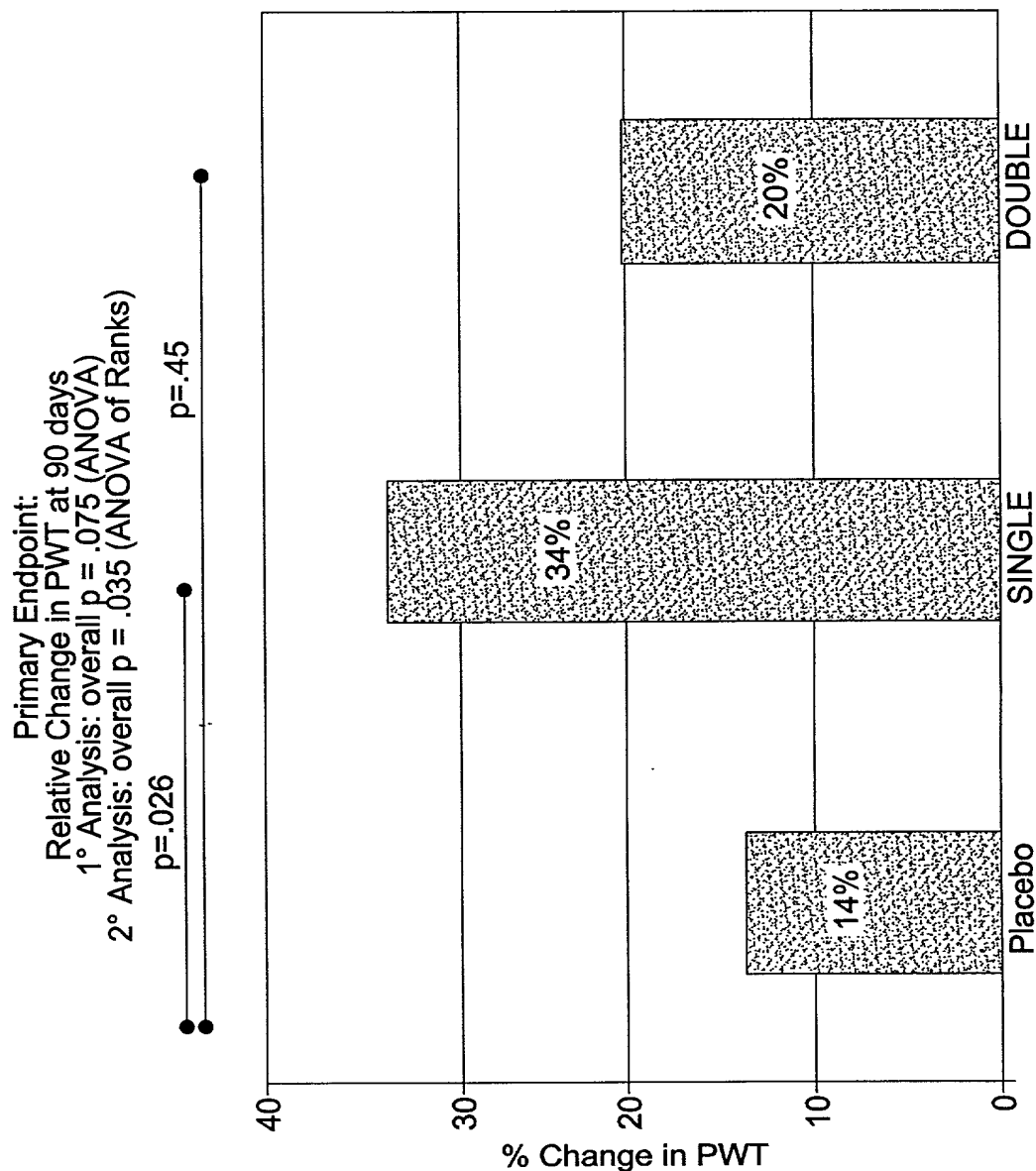


FIG. 6.

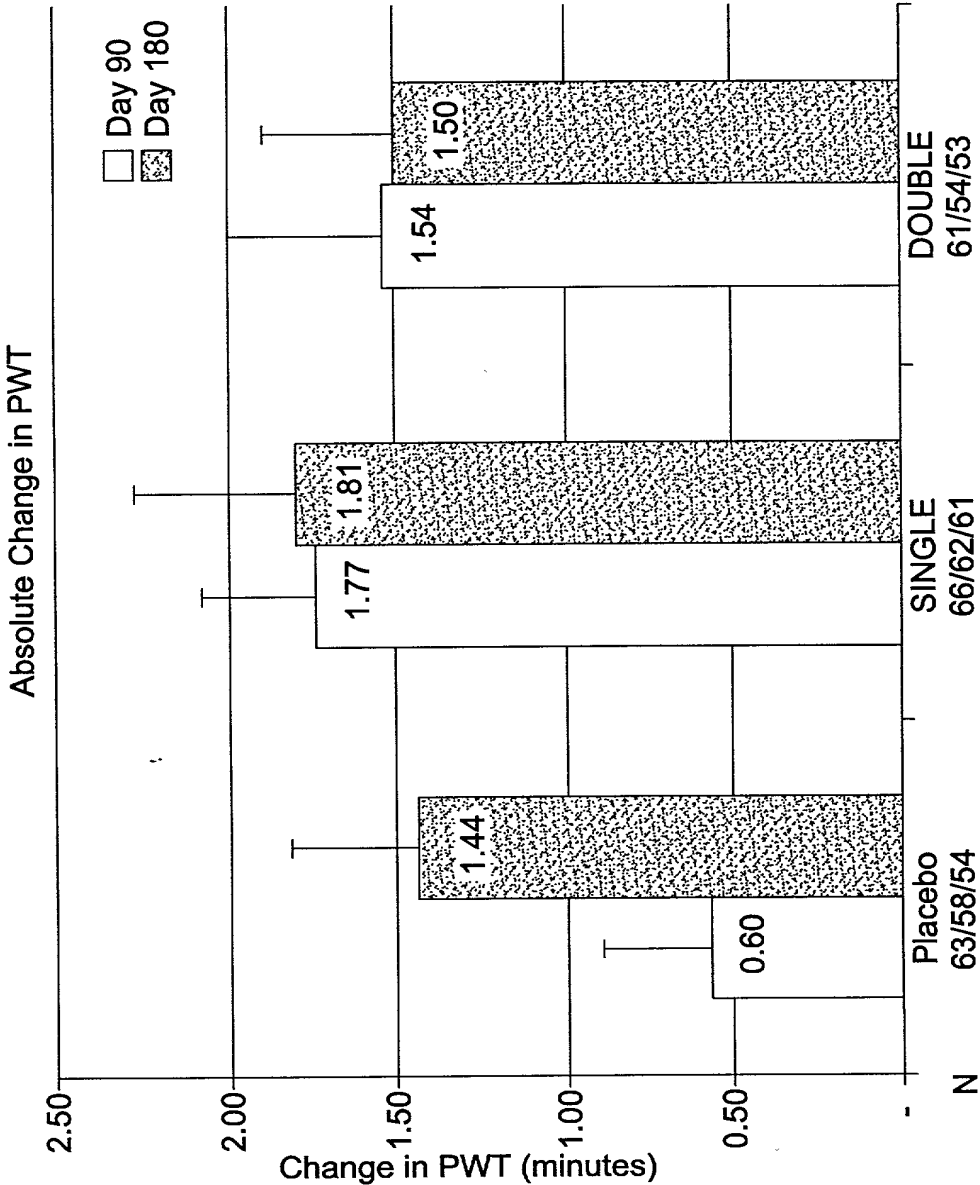


FIG. 7.

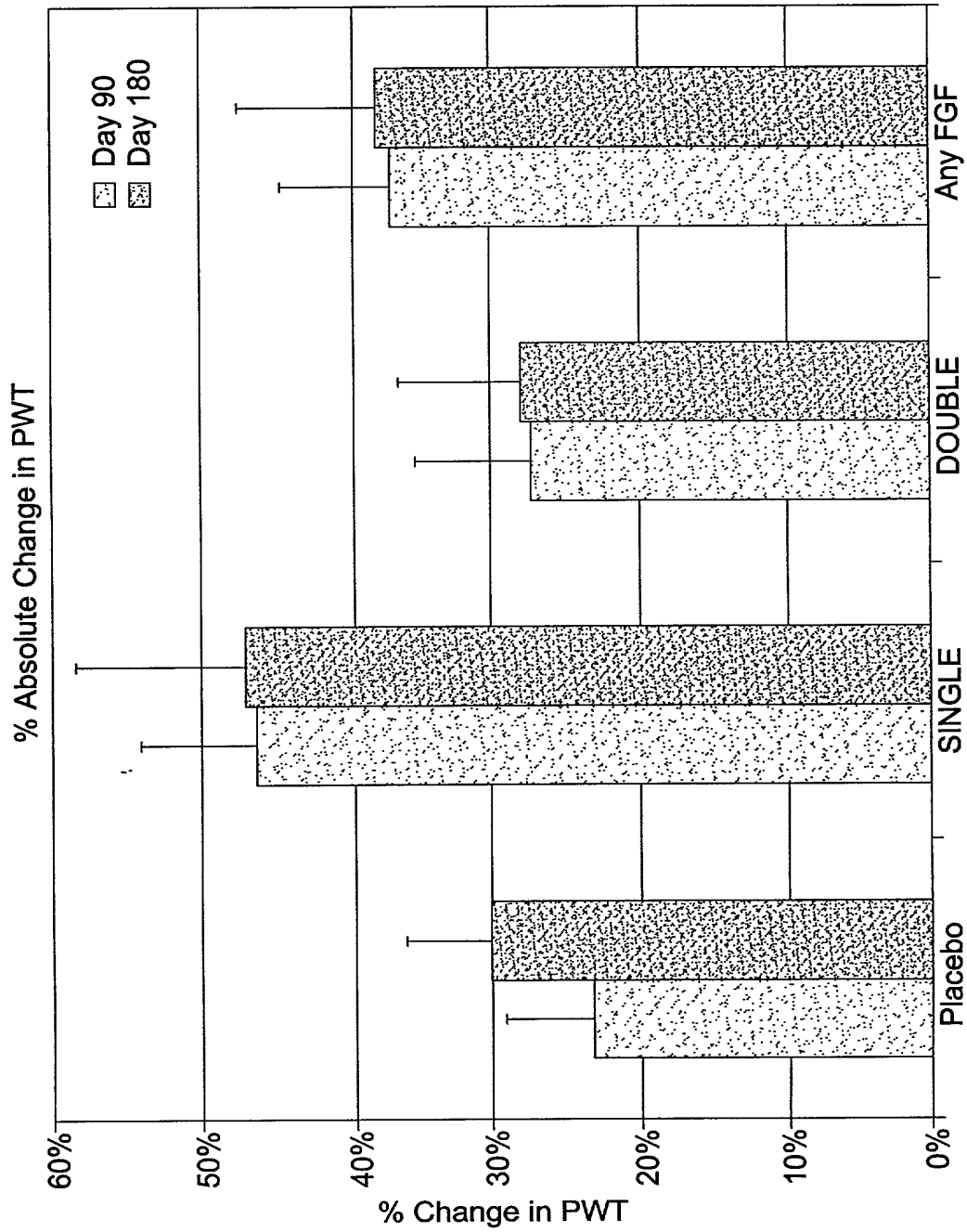
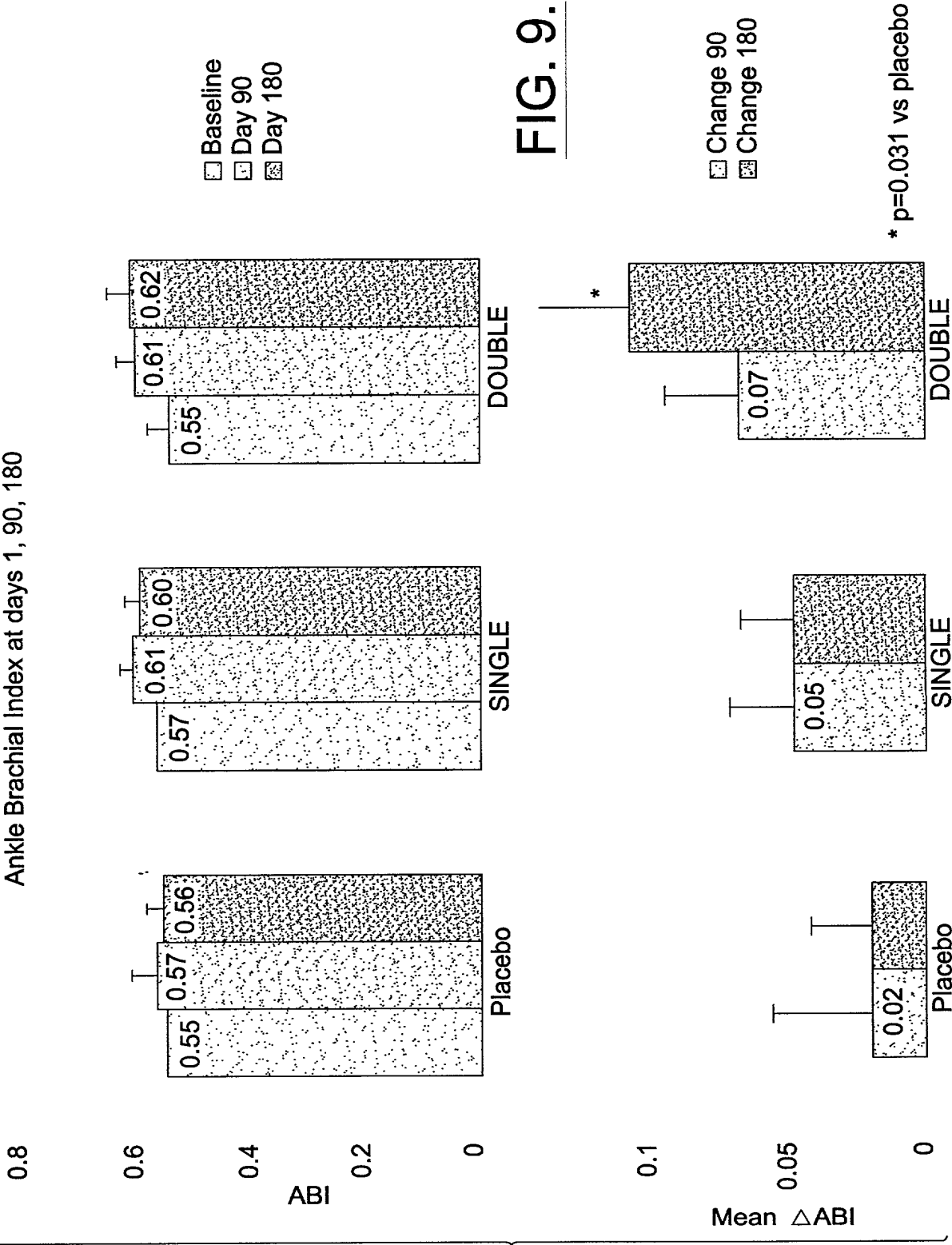


FIG. 8.

FIG. 8



10/20

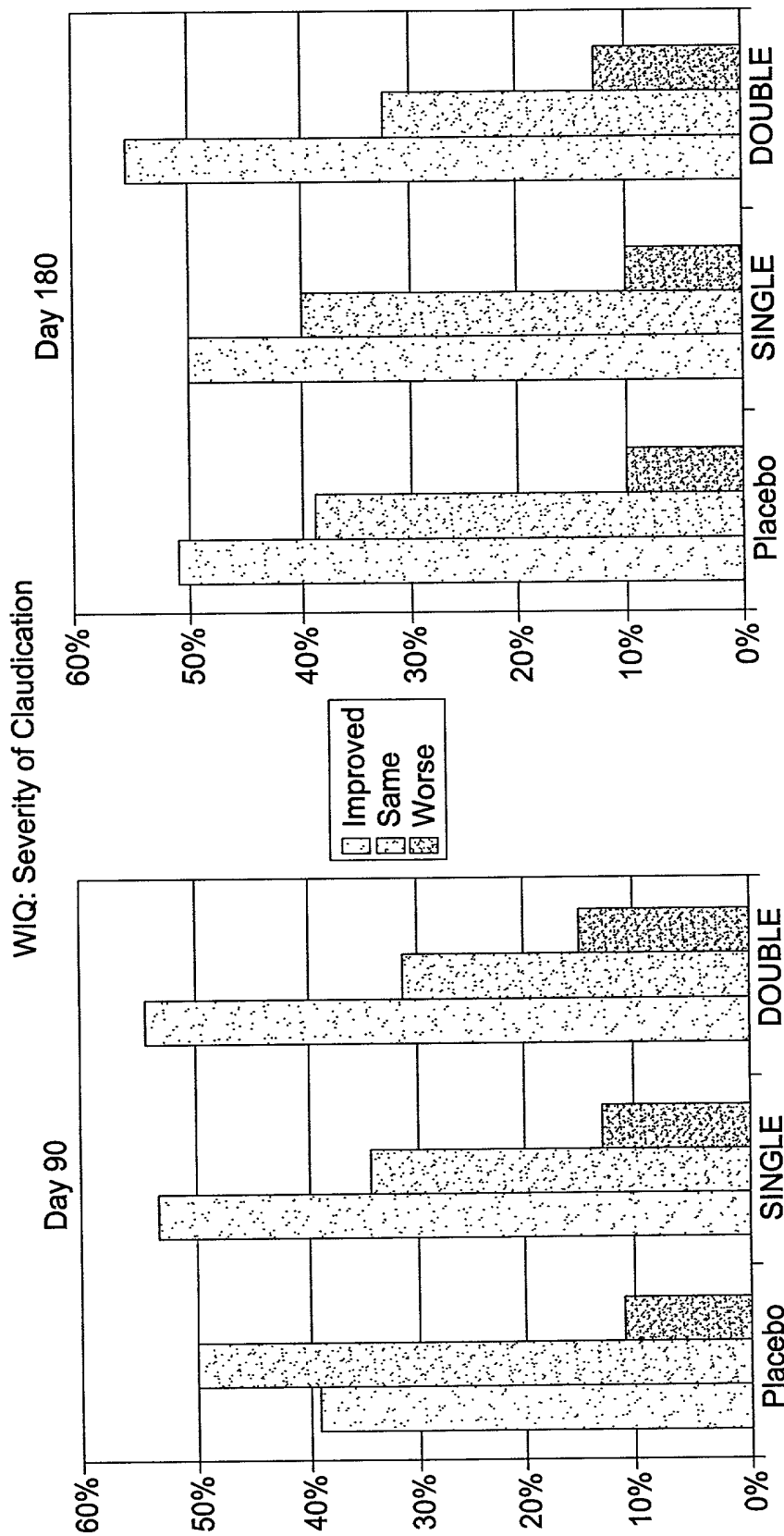


FIG. 10.

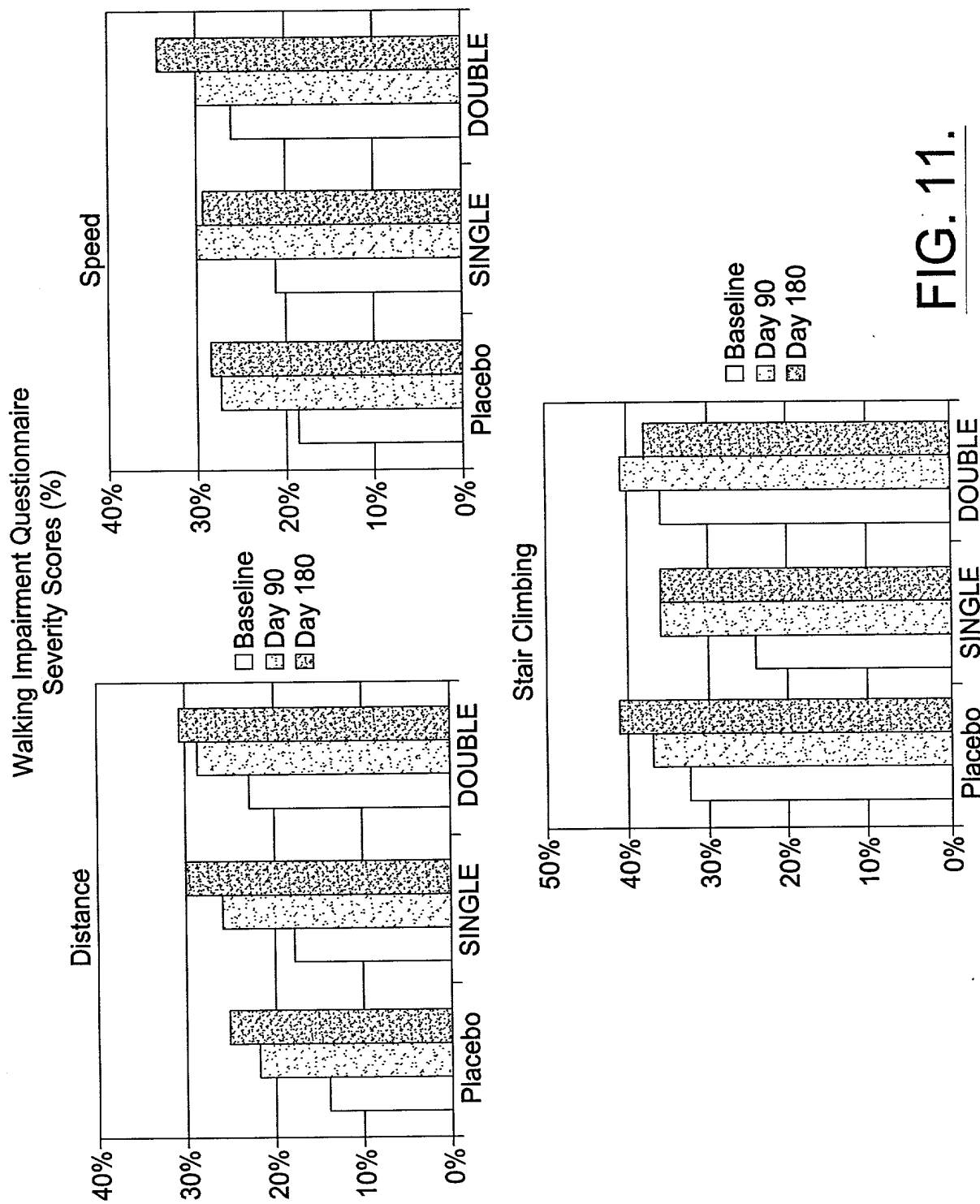


FIG. 11.

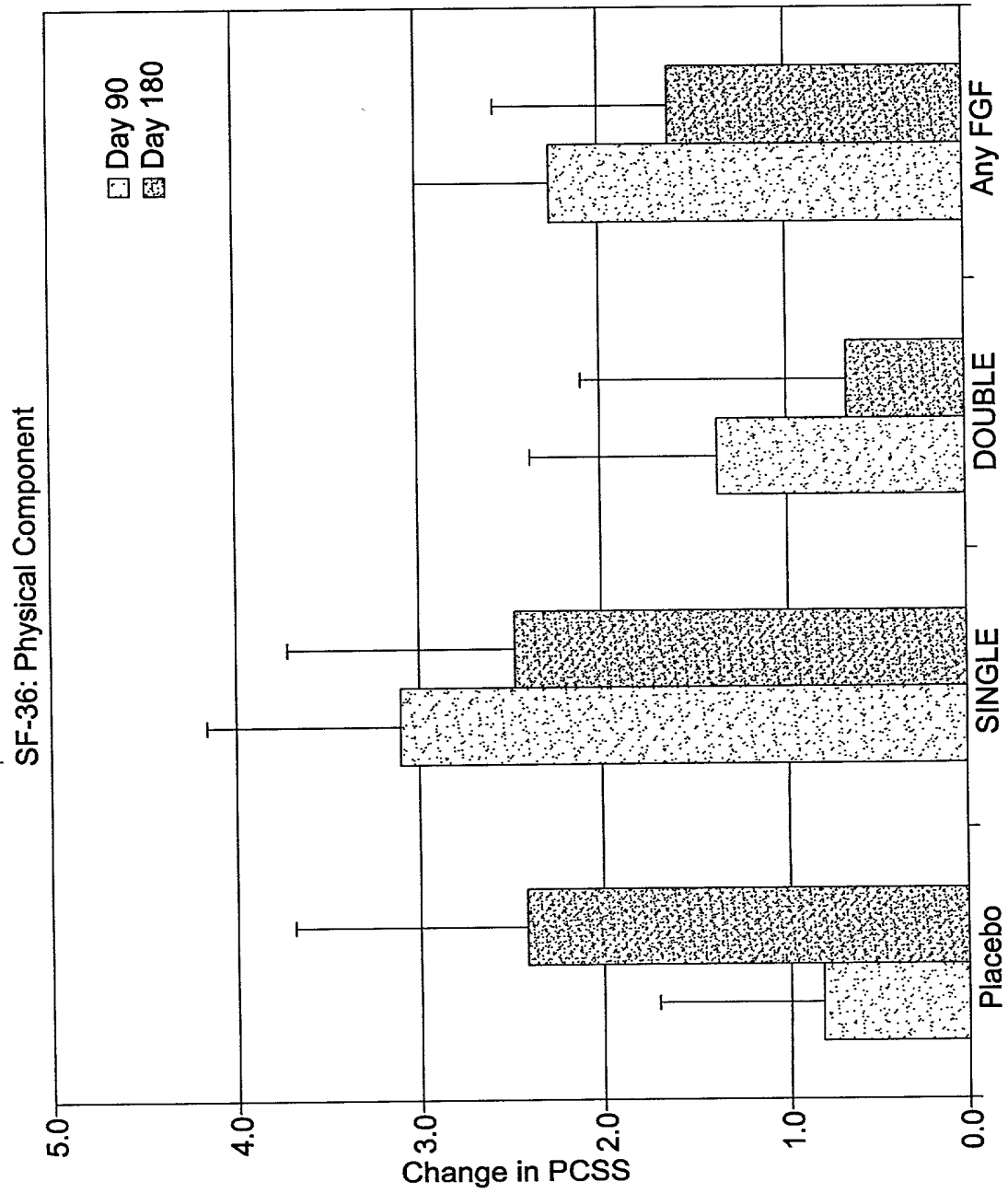


FIG. 12.

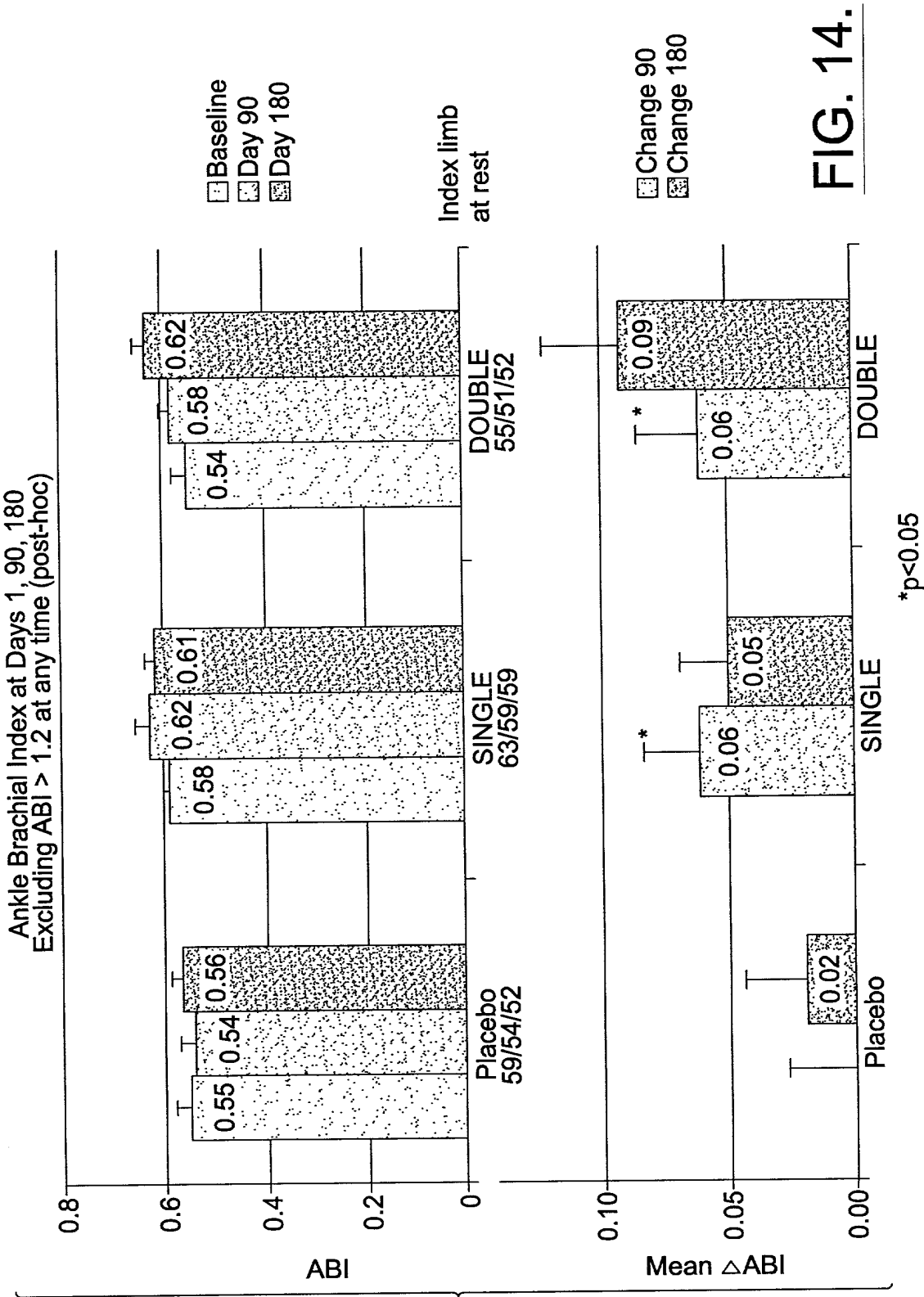
All PAD: Summary of Results

VARIABLE	Day 90	Day 180
PWT - primary overall	++	
PWT: pairwise/geometric	+++/+	+/-
Claudication Onset Time	+/+	-/-
Ankle Brachial Index	+/+	+/+
WIQ: claudication	+/+	-/+
WIQ: distance	0/-	+/-
WIQ: speed	+/-	+/0
WIQ: stairs	++/+	+/-
SF-36: Physical Score	++/+	+/-
SF-36: Mental Score	+/-	+/-

+++ = $p < .05$; ++ = $p < .15$; +, 0, - = directional change relative to placebo
SINGLE/DOUBLE

FIG. 13.

0986856-100501



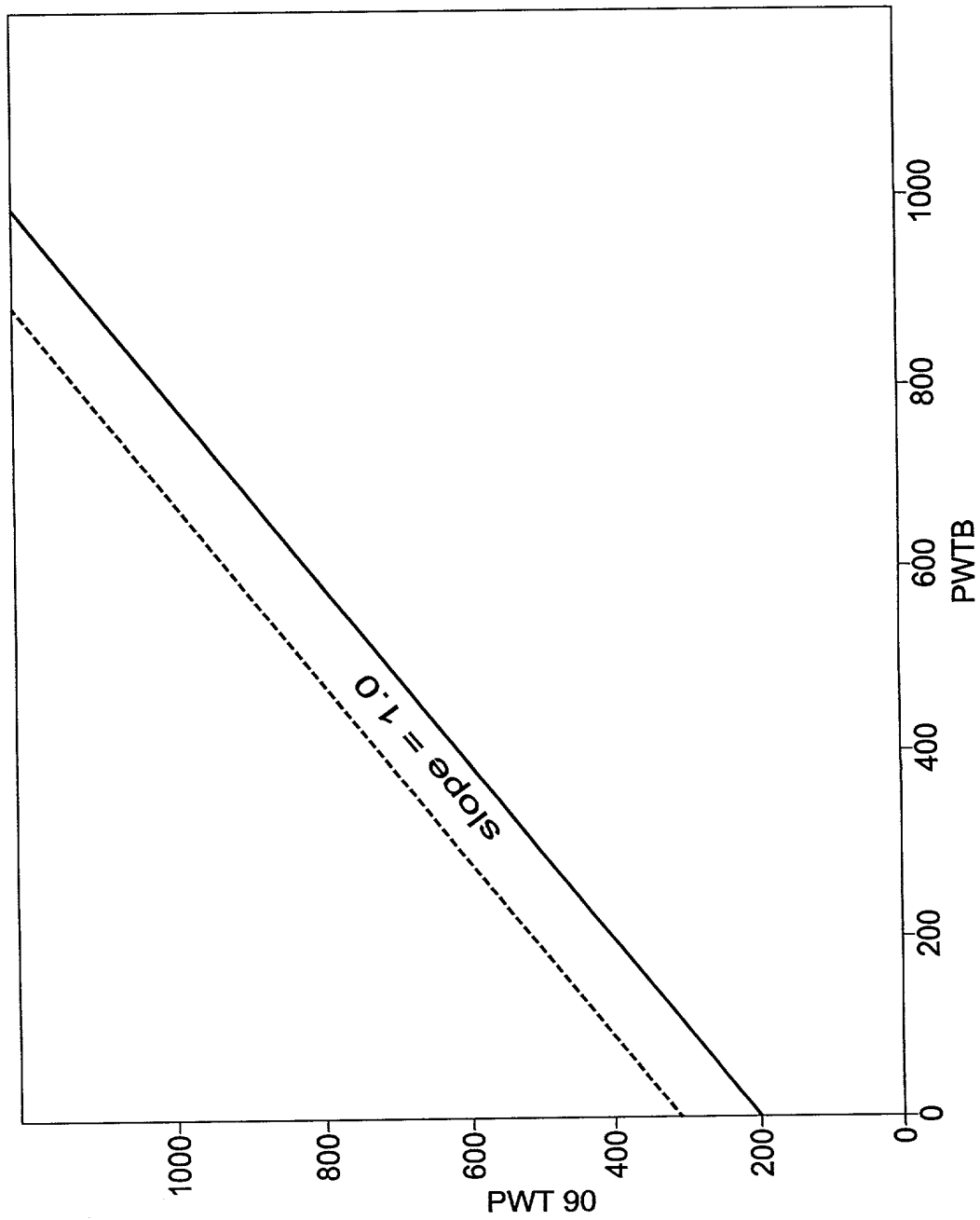


FIG. 15.

09886856 105001 95898860

Title: Methods and Compositions for the Treatment of Peripheral
Artery Disease

Inventor(s): Martha Jo Whitehouse

Application No: 09/886,856

Atty Dkt No: PP16090.004 (35784/235886)

16/20

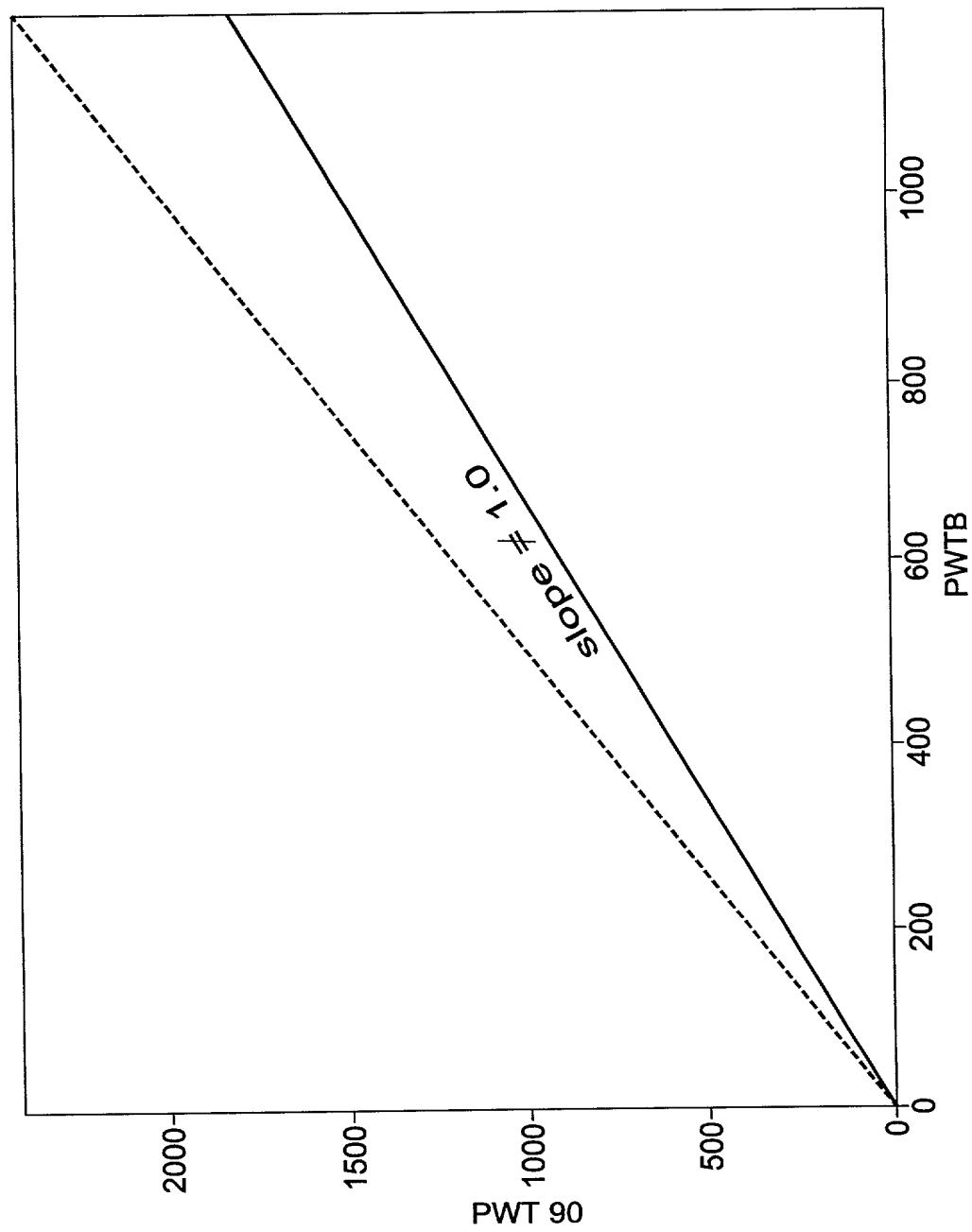


FIG. 16.

FIG. 16

Title: Methods and Compositions for the Treatment of Peripheral Artery Disease

Inventor(s): Martha Jo Whitehouse

Application No: 09/886,856

Atty Dkt No: PP16090.004 (35784/235886)

17/20

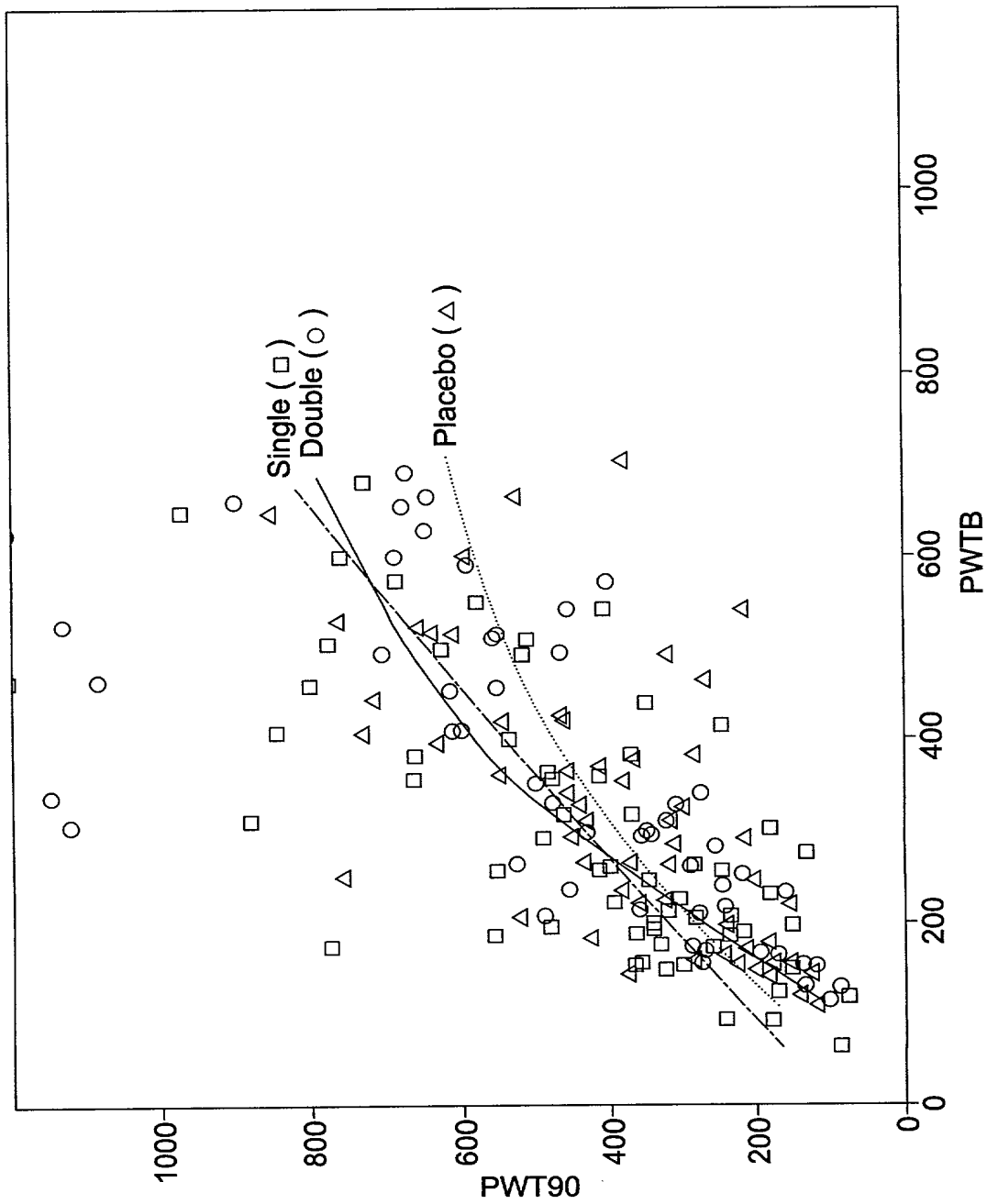


FIG. 17.

Title: Methods and Compositions for the Treatment of Peripheral
Artery Disease

Inventor(s): Martha Jo Whitehouse

Application No: 09/886,856

Atty Dkt No: PP16090.004 (35784/235886)

18/20

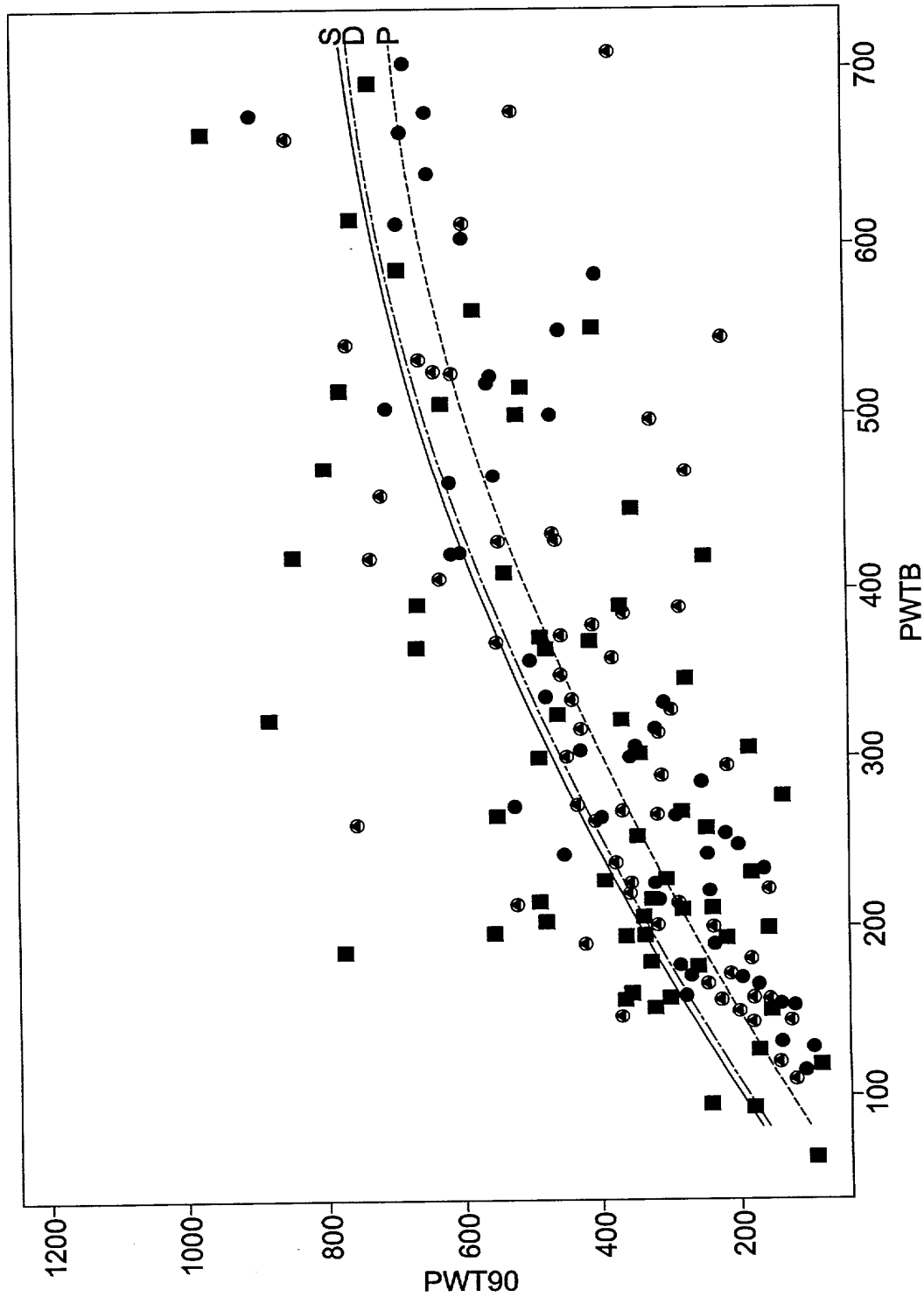


FIG. 18.

FIG. 18

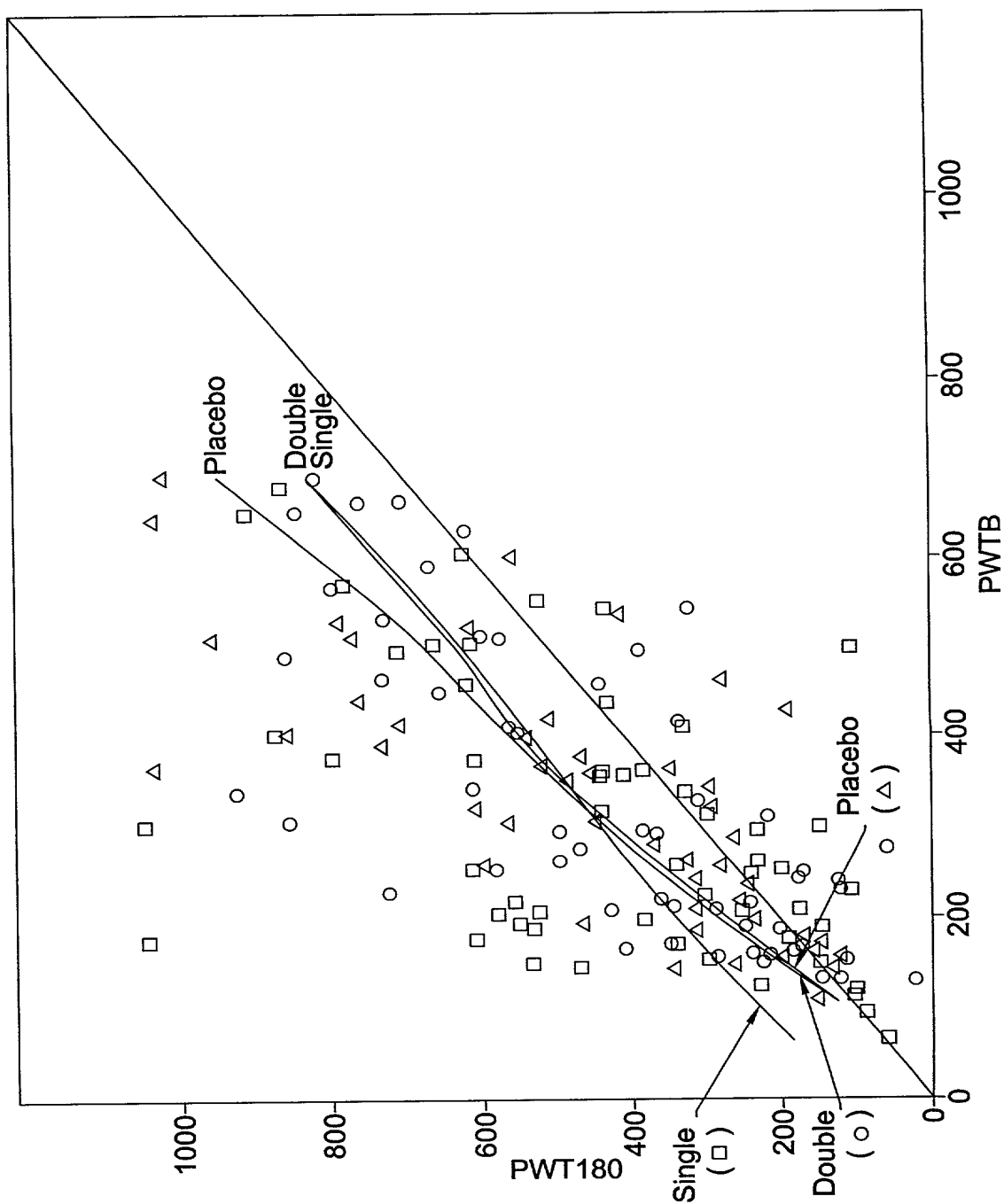


FIG. 19.

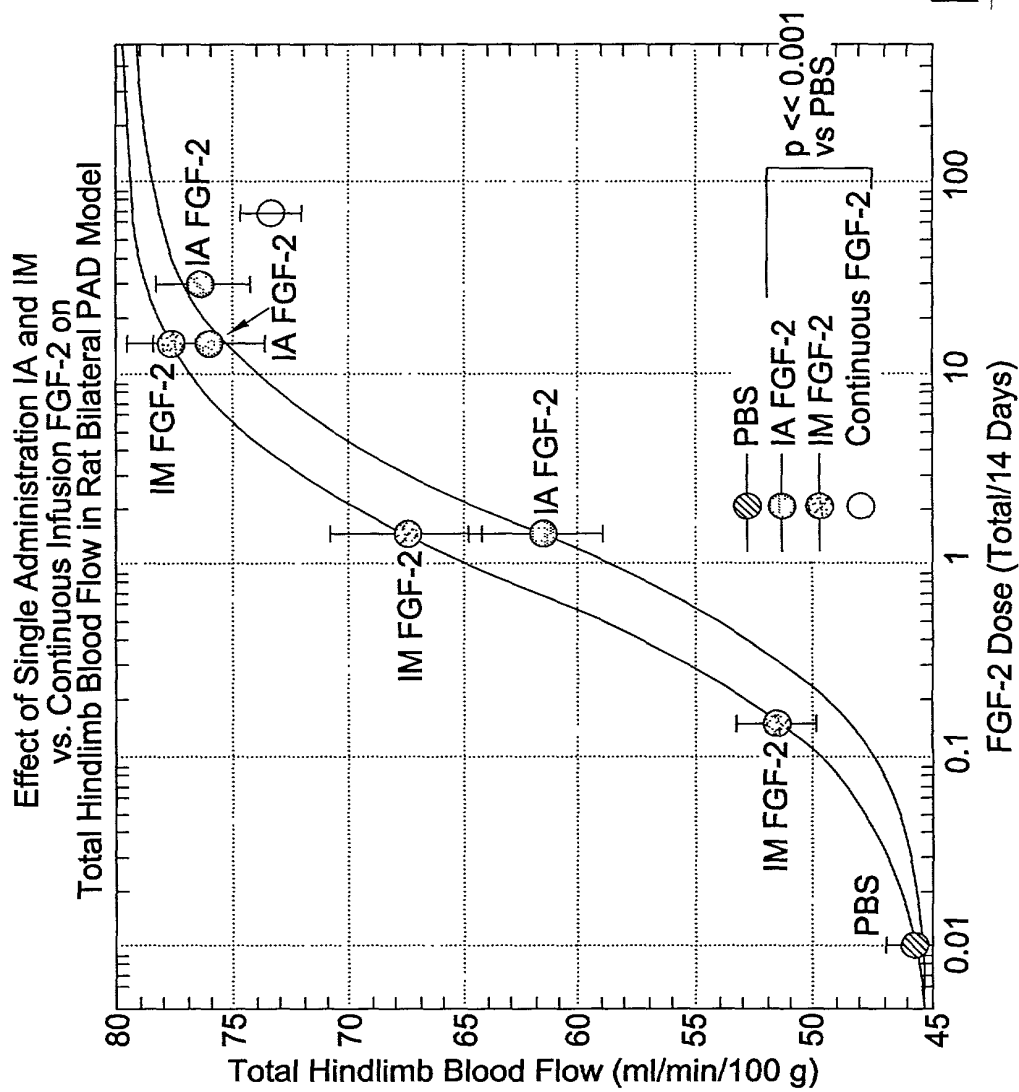


FIG. 20.